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Result

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Score

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SUMMARIES

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is the number of results predicted by chance to have a ater than or equal to the score of the result being printed, rived by analysis of the total score distribution.
                                                                                                                                                                                                            ALIGOTO Drosophil
BL281 T24Dl1-56
ALIGOTO Drosophil
BL2981 T24Dl1-56
ALIGOTO Drosophil
AZIJ3646 RPCI-23-4
Z29262 F. Tubripes
ALIJ3619 F. Iubripes
ALIJ3619 F. Iubripes
ALIJ66732 Drosophil
Z2071 F. Tubripes
ALIJ66741 F. Tubripes
ALIJ66743 REBAUBTO
AMIJ6318 NIOA425e
BEZ33928 EST403677
ALIJ6811 Drosophil
ALIGOTO Tetradon
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AL073856 Drosophil
AL073856 Drosophil
AL011359 F.rubripe
AL103838 Drosophil
AL300850 Tetraodon
AL063921 Drosophil
AW329517 N200762e
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AL108721 Drosophil
AL104216 Drosophil
Bl3042 T30M24-Sp6.
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AL169549 Tetraodon
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AUTHORS
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NNGGANNNANNNAANATGTTRCAATWAWAAGTATRGAAAAAAATAARWAAAAAA----
                                                                                                                                    WTGAWAAAGAAAAAAATAATATATWTAAAATAAATTAGWAGRATARATAAWTGAAATAA
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                                                 gaacactaaaattacaactttcagctgctgcaaatgaagatacagtaaacgtaaatactg
                                                                                                         AAAARAARAWAWAAAAAAGARAAAAANNNNNNNANNANNNNNAAANNNNAANNNN
                                                                                                                                                                                                                    <u>AAGAAARAAAAAAAAATWAAAWRAAAAACATAATRWGWGTAAAAACAAAAAAAAAGAAAAA</u>
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Determination of this BAC-end sequence was carried out as part of a collaboration with the European Drosophila Genome Project (EDGP) - collaboration with the European Drosophila melanogaster BAC http://www.edgp.ebl.ac.uk - This Drosophila melanogaster BAC library (Dros BAC) was made by Alain Billaud at CEPH (Centre d'Etude du Polymorphisme Humain) with funding provided by a MRC project grant. The DNA was prepared from embryos by Alain Bucheton and Genevieve Payan. It has been constructed in the vector
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Drosophila melanogaster genome sur
BACN9507 of DrosBAC library from
fly), genomic survey sequence.
AL102403
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Eukaryota; Metazoa; Arthropoda; Tracheata; Hexapoda; Ins
Eukaryota; Neoptera; Endopterygota; Diptera; Brachycera;
Pterygota; Neoptera; Endopterygota; Diptera; Brachycera;
Muscomorpha; Ephydroidea; Drosophilidae; Drosophila.
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Pred. No. 0.
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RCCarter, J., Clifton, S., Marra, M., Hillier, L., Kucaba, T., Martin, J., Beck, C., Wylie, T., Underwood, K., Steptoe, M., Theising, B., Allen, M., Bowers, Y., Person, B., Swaller, T., Gibbons, M., Pape, D., Harvey, N., Schurk, R., Ritter, E., Kohn, S., Shin, T., Jackson, Y., Cardenas, M., McCann, R., Waterston, R. and Wilson, R.

The Washington Univ. Nematode EST Project, 1999
Unpublished (1999)
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The Washington Univ. Nematode EST Project, 1999
Washington University School of Medicine
4444 Forest Park Parkway, Box 8501, St. Louis, 1
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BE224646
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Fax: 314 286 1810
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                                                                                                                                                           /note-"Vector: Lambda Uni-ZAP KR (Stratagene); Site 1: ECORI; Site_2: XhoI; mRNA was purified from 4 x 10E5 filariform larvae which had been Isolated from infected humans. CDNA was constructed and, using adaptors, was cloned unidirectionally into the vector from the ECORI site to the XhoI site. The library has an unamplified titer of 1.5 x 10E6 pfu/ml and an amplified, undiluted titer of 7 x 10E9 pfu/ml. The average insert size of the unamplified library is 975 bp (range, 500-1500)."
                                                                                                                                                                                                                                                                                                                                                                /strain="Filariform larvae obtained from humans"
/db_xref="taxon:6248"
/clone_lib="TBN95TM-SSFH"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Web: www.genoscope.cns.fr)

Determination of this BAC-end sequence was carried out as part of a collaboration with the European Drosophila Genome Project (EDGP) - http://www.edgp.ebi.ac.uk - This Drosophila melanogaster BAC http://www.edgp.ebi.ac.uk - This Drosophila melanogaster BAC library (Dros BAC) was made by Alain Billaud at CEPH (Centre d'Etude du Polymorphisme Humain) with funding provided by a MRC project grant. The DNA was prepared from embryos by Alain Bucheton
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/plasmid="pseloBaC11"
/db_xref="taxon:7227"
/clone_lib="DrosBaC"
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6; Mismatches 365;
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AL169549.1
GSS; genome
                                                                                                                          Submitted (12-APR-2000) to the EMBL/GenBank/DDBJ databases This sequence is a single read and was generated as part of scale cione-end sequencing project of the Tetraodon nigrovi genome. For more information, please take a look at http://www.genoscope.cns.fr/Tetraodon.
                                                                                                                                                                                                                                                                                                                                           2 (bases 1 to 773)
Roest-Crollius, H., Jaillon, O., Dasilva, C., Bouneau, L., Fisher, C. Bernot, A., Fizames, C., Wincker, P., Brottler, P., Quetler, F., Saurin, W. and Weissenbach, J.
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Roest-Crollius, H., Jaillon, O.,
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   ы
                                                     /organism="Tetraodon nigroviridis"
/db_xref="taxon:99883"
/clone="199024"
/clone_11b="G"
/note="Genoscope sequence
265 c 69 g 184
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er,F., Saurin,W., Bernot,A.
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                                   Roest-Crollius, H., Jaillon, O., Bernot, A., Fizames, C., Wincker, Saurin, W. and Weissenbach, J.
                                                                                                                                                                                      1 (bases 1 to 735)
Roest-Crollius, H., Jaillon, O., Dasilva, C., Fizames, C., Fisher,
Bouneau, L., Billault, A., Quetier, F., Saurin, W., Bernot, A. and
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                  Human gene number estimate
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Tetraodontiformes; Tetraodontoidei; Tetraodontidae; Tetraodon.
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r,P., Quetier,F.,
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Submitted (23-JUL-1999) Genoscope -
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BP 191 91006 EVRY cedex - FRANC
                                                                                                                                                                                                                                                                                            fly), genom:
AL108447
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GSS:
                                                                                                                                     Drosophila melanogaster
Eukaryota; Metazoa; Arthropoda; Tracheata; Hexapoda; Insecta;
Pterygota; Neoptera; Eudopterygota; Diptera; Brachycera;
Muscomorpha; Ephydroidea; Drosophilidae; Drosophila.

1 (bases 1 to 917)
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                                                                                                                           Genoscope.
                                                                                                                                                                                                                                                                        fruit fly.
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nilarity 45.7%;
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/db_xref="taxon:99883"
/glone="123M05"
/clone_lib="g"
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Pred. No. 0.016;
5; Mismatches 194;
                                              a - Centre National de sequencage
  (E-mail : seqref@genoscope.cns.fr
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GSS; genome survey sequence.
GSS; genome nigroviridis.
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Tetraodon nigroviridis
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Eukaryota; Metazoa; Chordata; Teleostei; Euteleostei; Neoteleostei;
Eutrypterygii; Chenosquamata; Acanthomorpha; Euacanthomorpha;
Holacanthopterygii; Acanthopterygii; Percomorpha;
Tetraodontiformes; Tetraodontoidei; Tetraodontidae; Tetraodon.
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freshwater pufferfish Unpublished 2 (bases 1 to 1076)
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                                                                                      l (bases 1 to 1076)
Roest-Crollius,H., Jaillon,O.,
Bouncau, E., Billault,A., Queti-
Weissenbach,J.
                                        Charaterization and repeat analysis of the compact freshwater pufferfish Tetraodon nigroviridis
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/db_xref="taxon:7227"
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/clone="BACN37E22"
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hes 116; Conservative
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Submitted (02-UN-1999) Genoscope · Centre National de Sequencage : BP 191 91006 EVRY cedex · FRANCE (E-mail : seqref@genoscope.cns.fr)

Web : www.genoscope.cns.fr)
Determination of this Back-end sequence was carried out as part of a coilaboration with the Berkeley Drosophila Genome Project (BDGP).

The BDGP is constructing a physical map of the Drosophila melanogaster genome using these BACs. For further information please see http://www.fruitfly.org The BDGP Drosophila melanogaster BAC ilbrary was prepared by Kazutoyo Osoegawa and haron Mammoser in Pieter de Jong's laboratory in the Department of
                                                                                                                                                                                                                                                                                                                                                                                                                                                               CNSOOCI3 560 bp DNA GSS 04-JUN-1999 DIGGOODHIA melanogaster genome survey sequence T7 end of BAC * BACR2505 of RPCI-98 ilbrary from Drosophila melanogaster (fruit fix), genomic survey sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Roest-Croilius, H., Jaillon, O., Dasilva, C., Bouneau, L., Fisher, C., Bernot, A., Fizames, C., Wincker, P., Brottler, P., Quetler, F., Saurin, W. and Weissenbach, J.

Human gene number estimate provided by genome wide analysis using Tetracodon nigroviridis DNA sequence
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rterygota; Neoptera; Endopterygota; Diptera; Brachycera;
Muscomorpha: Ephydroidea; Drosophilidae; Drosophila.
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                             Direct Submission
Submitted (23-JUL-1999) Genoscope - Centre
BP 191 91006 EVRY cedex - FRANCE (E-mail:
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                                                                                                                                                                                                                                      Drosophila melanogaster genome survey sequence SP6 end of BACN37807 of DrosBAC library from Drosophila melanogaster fly), genomic survey sequence.
ALL108721
   Determination
                                                                                    Genoscope:
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/db_xref="taxon:7227"
./cione=lib="RPCI-98"
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www.genoscope.cns
nation of this BAC
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AL104216
AL104216.1 GI
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                       Eukaryota; Metazoa; Arthropoda; Tracheata; Hexapoda; Insecta; Pterygota; Neoptera; Endopterygota; Diptera; Brachycera; Muscomorpha; Ephydroidea; Drosophiiidae; Drosophiia.

(bases 1 to 1025)
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Drosophila melanogaster genome survey sequence T7 end of BAC
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and Genevieve
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gcaaatgaagataca
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Determination of this BAC-end sequence was carried out as part of a collaboration with the European Drosophila Genome Project (EDGP) - http://www.edgp.ebi.ac.uk - This Drosophila melanogaster BAC library (Dros BAC) was made by Alain Billaud at CEPH (Centre d'Etude du Polymorphisme Humain) with funding provided by a MRC project grant. The DNA was prepared from embryos by Alain Bucheton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         submitted (23-JUL-1999) Genoscope - Centre National de Sequencage
BP 191 91006 EVRY cedex - FRANCE (E-mail : seqref@genoscope.cns.fi
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Arabidopsis Thaliana Genome Center
University of Pennsylvania
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BAC End Sequences at ATGC Unpublished (1997) Other_GSSs: T30M24-Sp6, T30M24-Sp6.2
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T30M24-Sp6.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Email: jecker@atgenome.bio.upenn.edu seq primer: sp6
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Fax: 215-898-8780
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l (bases 1 to 1147)
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/note="Yector: BeloBACI; Site_1: HindIII;
/Produced by Rod Wing"
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553 t 541 Others
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/strain="Columbia"
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              and how to order individual BAC clones, the entire library, of filters for hybridization from the BACPAC Resource Center can found at http://bacpac.med.buffalo.edu/drosophila_bac.htm.
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Web: www.genoscope.cns.fr)
Determination of this BAC-end sequence was collaboration with the Berkeley Drosophila the BDGP is constructing a physical map of
                                          Direct Submission
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- Web : www.genoscope.cus.fr)
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                                                                                                                                                                                   CNSOOHX9. 1101 bp DNA GSS 03-JUN-1999
Drosophila melanogaster genome survey sequence TET3 end of BAC:
BACR35M04 of RPCI-98 llbrary from Drosophila melanogaster (fruit fly), genomic survey sequence.
ALD73856
Genoscope.
Direct Submission
                                                      Drosophila melanogaster
Eukaryota; Metazoa; Arthropoda; Tracheata; Hexapoda; Ins
Pterygota; Neoptera; Endopterygota; Diptera; Brachycera;
Muscomorpha; Ephydroidea; Drosophilidae; Drosophila.
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/db_xref="taxon:7227"
/db_xref="RPCI-98"
/clone="BACR36C06"
/note="end; TET3"
133 c 93 g I50 t 91
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                       tgtaagtgcgattaacactcaaaacaaagctgttgaattaacagcagtaccagtgaacgg 857
                                                                       agaagtagcgaaaaaagcattagatgcagatgaagctgcgcttactccaaaagttgaaag
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/db_xref="taxon:7227"
/clone_lib="RPCI-98"
/clone="BACR35M04"
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Fugu rubripes.
Fugu rubripes
Eukaryota, Metazoa; Chordata; Vertebrata; Actinopterygil;
Eukaryota, Metazoa; Chordata; Vertebrata; Actinopterygil; Per
Neopterygil; Teleostei; Euteleostei; Acanthopterygil; Per
Tetraodontiformes; Tetraodontoidei; Tetraodontidae; Fugu.
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Elgar,G., Clark,M., Smith,S., Meek,S., Warner,S., Umrania,Y.,
Williams,G. and Brenner,S.
Direct Submission
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./db_xref="taxon:31033"
/clone=11b="cosmid 016E10"
./clone="016E10ac6"
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